

Sequence Listing

SEQ ID NO.: 1: ESX cDNA ORF and deduced amino acid sequence
(See Figure 1)

SEQ ID NO.: 2: ESX cDNA sequence (5' untranslated + ORF + 3' untranslated) 1907 b.p.

cgccagatacctcagcgctacctggcggaactggatttctctcccgccctgccggcctgcct
gccacagccggactccgccactccggtagcctcatggctgcaacctgtgagattagcaacat
tttagcaactacttcagtgcgatgtacagctcggaggactccaccctggcctctgttcccc
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gatgctgggacccttaggatggggctcccagctccttctcctgtgaatggaggcagagacc
tccaataaagtgccttctgggctttttctaaaaaaaaaaaaaaaaaaaaa

SEQ ID NO.: 3: Complete ESX deduced amino acid sequence (see Figure 1)

SEQ ID NO.: 4: First variable region (nucleotides 1-189 of Figure 1)

atggctgcaacctgtgagattagcaacatttttagcaactacttcagtgcgatgtacagctc
ggaggactccaccctggcctctgttccccctgctgccacctttggggccgatgacttggtag
tgaccctgagcaacccccagatgtcattggagggtacagagaaggccagctgggttgggggaa
cag

SEQ ID NO.: 5: ❖ Pointed❖ region (nucleotides 190-309 of Figure 1)

ccccagttctggctcgaagacgcaggttctggactggatcagctaccaagtggagaagaacaa
gtacgacgcaagcgccattgacttctcacgatgtgacatggatggcgccaccctctgc

08978217.112597

SEQ ID NO.: 6: Second variable region (nucleotides 210-561 of Figure 1)

aattgtgcccttgaggagctgcgtctggtcttttgggcctctgggggaccaactccatgcccc
gctgcgagacctcacttccagctcttctgatgagctcagttggatcattgagctgctggaga
aggatggcatggccttccaggaggccctagacccagggccctttgaccagggcagccccctt
gccaggagctgctggacgacggtcagcaagccagcccctaccaccccggcagctgtggcgc
agga

SEQ ID NO.: 7: Deduced amino acid sequence for second variable region (amino acids 104-187 of Figure 1)

asn-cys-ala gly-ala-gly

SEQ ID NO.: 8: Serine-rich region (nucleotides 562-714 of Figure 1)

gccccctcccctggcagctctgacgtctccaccgcagggaactgggtgcttctcggagctccca
ctcctcagactccggtggaagtgacgtggacctggatcccactgatggcaagctcttcccca
cgcgatgggttttcgtgactgcaagaagggg

SEQ ID NO.: 9: Third variable region (nucleotides 715-819 of Figure 1)

gatcccaagcacgggaagcggaaacgaggccggcccccgaagctgagcaaagagtactggga
ctgtctcgagggcaagaagagcaagcacgcgcccagaggcacc

SEQ ID NO.: 10: Ets DNA Binding domain (nucleotides 820-1062 of Figure 1)

cacctgtgggagttcatccgggacatcctcatccacccggagctcaacgagggcctcatgaa
gtgggagaatcggcataagggcgtcttcaagttcctgcgctccgaggctgtggcccaactat
ggggccaaaagaaaaagaacagcaacatgacctacgagaagctgagccgggccatgaggtac
tactacaaacgggagatcctggaacgggtggatggccggcgactcgtctacaagttt

SEQ ID NO.: 11: Fourth variable region (nucleotides 1063-1113 of Figure 1)

ggcaaaaactcaagcggctggaaggaggaagaggttctccagagtcggaac

SEQ ID NO.: 12: C-terminal 16 amino acids (amino acids 356-371 of Figure 1)

lys-asn-ser ser-arg-asn

SEQ ID NO: 13 5'ESX-DBD

5'-CCGGGACATCCTCA TCCACCC-3'

SEQ ID NO: 14 3' ESX-DBD

5'-GTACCTCATGGCCCGGCTCAG-3' (SEQ ID NO. 14)).

SEQ ID NO: 15 Mouse ESX genomic sequence.

1 GGATCCTTCC AAGGCACTGA CCTCACCCAA TTCTTTCTCA CTTTCTCCT
51 CCATTAACT GTGGACGGAA TCAATACTCA GGGGGATGCG CTAGCTCTAA
101 GATTTCTGCA GCTTTGCCTC TCCTGAGCGG AAGCCCCGTG AAGGCAAGGG
151 AGCTAGCTGA TGGACTCTTT GTGGTCTTCT TCCTCTTTGC TCTGGAGACC
201 CAACCAGGTG TTCTTAGGGG AAGGAGCACG TGAGTAGCCA AGAGGCTAAA
251 AGCTGGTTCT CCCACATTCC AGGGTAAGTG ACTGGGTAGA GGGTGTGTCT
301 GCCTCAGGCT GCTTGGAGGA GGTCCCCTGA AGGGCCATGA GAAAATCCTA
351 CCCAGAGCCC TTGGTTTTCC AGCAGCCCTC CACCTAGAGG AAAGGAGCCT
401 GTCGTTCTGA AGATGAAGAG TGGAGCCTAT GGGGGTGGGC AGATTGTGTC
451 CTGGGACAAT GGGGTACCTA GAAGAGAAAG GAATCTCCTT TCGTTTGAGG
501 TCTACCTGGG GGTCTGTGTG CTGTAAATGG GGTGGAGAGA GGAGAAGACA
551 CAGATCTTAT AACGTAGATG CAGGAAATGC TGACAGTTCA GTGTAGAGAA
601 CTTACTCAAT TCATATAGCC TCCAAAGCTA TCTCCTCAGG CAACGCAAAA
651 CAAACCAGTT GGAGCCGCAA GACATCTAAT GGCTTATCGA GTCCACACC
701 CTCGATTCTT TGCTAATTTT ATGGTTTTGC TTTTGAGACA ATCTACTGTA
751 GCCTAAGATA GCCCCAACT CAAATGTAGC TGAGGCTGAC TGACCCTGAG
801 CTCTGGAATT CCAGACACAT GCATATCTTT TGCTAGGCAA TAATCGCTCT
851 ACCAGCTGTA CTCCCACATT CCAGGGTAAG TGA CTGGAAT TCTCACTTAC
901 TATATCCCTT TAAAAATTCC CTGAGTGGGA TGGTTGTAGC CAGAGGGAAA
951 AGGCACCAAC AACTGCTTGT CACTTTCCTA ATTTGGTAGC CTGAACAAAC
1001 CACTTATCAA GACAACAAC ATATATCATT TCTTTTCTTC TCTCTCTCTC
1051 TCTCTCTCTC TCTCTCTCTC TCTCTCTCTC TCTCTCTCTC TCTCTCTTTN
1101 GAAAGAGTCT CACTACTATG TAGCCCTTGA TAACCTAGAA CTCACTATGT
1151 AGTCCAGGCT TGGCCTTCAG CTCGCAGAGG TCCACTTGCC TTGGGAGTTG
1201 AGAGATTAAA GGGATGCATC TCCACATGTG TCCAACAGTG CTTTTTAAAA

1251 ATATTTTAA AACCATGCTT ACAGCCAGGC ATAGTGGGCG TGCCTTTAAT
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 1351 CACATAGTAA GTCCCAGGAT AGCTAGAACT ATGTAAAGAC CATGTCTCAA
 1401 AAAAGATGCA CACACACATA TACACACACA CGTTTGTATG TGTTTGTTTA
 1451 GTGTGTATGT GTGTGTACAC TTGCACATAA AGGTCAGAGT ACCACATTAC
 1501 AGGAGTCAGT TTTCTCCTTT TATCATGTAT GGATGGAACA CGGGTCCATC
 1551 CATAGCATCC TTAGCAGCAG GTATCCTTAT CCACTGAGCT ATCTCAGCAG
 1601 CCCACATTG CTTATTGGAT GTTTTTGGAT GAGGATAGTT ATATTAAAAA
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 1701 TGTCTCTTGA GTTTGAGGCC AGCCTGGTAT ATGTAGCTAG ACAAAGTTTC
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 1851 AGTTTTAGTT TTTCAAGACA GAGTTTCTCT GTGTAGCCCT GGCTGTCCTG
 1901 GAACTCACTC TGTAGACAAG GCTGGCCTCG AACTCAGAAA TCCTTCTACC
 1951 TCTACTTCAG GAGTGCTGGG ATTAAAGATG TGCGCTGCCC TCCTCCACCC
 2001 CAATTTGTTT TTGTTTTTTA AGGGCCCCGG TAAACAGTAA ATTAACATGT
 2051 GCATCCTGTT TGTCTTTGTA ATGACTCAA TGTTGGGCTT CTGACCACTA
 2101 GAGGGCAGCA GGCAGATACT AATGGACTGG GCGGAGAGAA GGGTAATCAG
 2151 GAGCAGACCA GACTCGCGGA TAAACCAAAC AGCACCGCCA GCCGACCCTA
 2201 GGCGAGGAGA GCGCCACAGG CACCAAGGGA AGACTTGAAG TAGTGTCTGA
 2251 TCTCTACCGC TTCAGCAACC ATCGCGTTTG GGTGGGCTCC AGACAGGCAA
 2301 AGTGCCAGCA AATGGTCCCT GTAGCTGACT AAACAGACTA TCAGACCCAA
 2351 ACCACCACTG GACCGTGAAT GTTGCCAGT GTGTTGCCTA GCCGCTTTCA
 2401 GAATCCCAGC TTCTGGGTGT TGTGGAGGAA ACCCCTTAGC CTCGGTAACT

2451 TTCACCAGGC CCTTCTTGTC TCTAGACATC TAGACAGTTG GAAGCATCAG
 2501 TCTTGACCCA GCCACCGGTT CAGATTCTTT GCCTTGCTTT TTCTTCCCA
 2551 GTTCAGCCCT GGCCAGGCCC CCAGGAAGAA TTTCCAGGGC CAGAGGGCAG
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 2651 CAGTACCCAG GGCCCAACCC CAGAGGGTGC GGAATGACAG ATTCTGACAA
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